WO 2005/077191 PCT/DK2005/000033

## SEQUENCE LISTING

| <110>  | Novozymes A/S                      |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
|--|------------------------------------|-----------------------|------------------------|------------------|-------------------|------------------|--------------------|------------------|------------------|-------------------|-------------------|-------------------|-----|--|
| <120>  | Preparation of Dough-Based Product |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <130>  | 10581-wo                           |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <160>  | 4                                  |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <170>  | PatentIn version 3.2               |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <210><br><211><br><212><br><213>                                       | DNA                                |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <220><br><221><br><222>  | > CDS                              |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <220> <221> sig_peptide <222> (58)(141)                                |                                    |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <220> <221> mat_peptide <222> (142)(687)                               |                                    |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   |     |  |
| <400> 1 aatcgacaac aaacgtgtaa ataagtagta cgataaaaat tttgaggagg acgaatc |                                    |                       |                        |                  |                   |                  |                    |                  |                  |                   |                   |                   | 57  |  |
| atg tt<br>Met Ph   | t aag ttc<br>e Lys Phe<br>-25      | gtt ad<br>Val Ti      | cg aaa<br>hr Lys       | gtt<br>Val       | ttg<br>Leu<br>-20 | acg<br>Thr       | gta<br>Val         | gta<br>Val       | att<br>Ile       | gca<br>Ala<br>-15 | gct<br>Ala        | aca<br>Thr        | 105 |  |
| att ag<br>Ile Se   | t ttt tgt<br>r Phe Cys<br>-10      | ttg ag<br>Leu S       | gt gca<br>er Ala       | gta<br>Val<br>-5 | ccg<br>Pro        | gca<br>Ala       | agt<br>Ser         | gct<br>Ala<br>-1 | Asn              | acc<br>Thr        | tat<br>Tyr        | tgg<br>Trp        | 153 |  |
| caa ta<br>Gln Ty<br>5  | t tgg acc<br>r Trp Thr             | gat g<br>Asp G<br>10  | ly Gly                 | gga<br>Gly       | aca<br>Thr        | gta<br>Val       | aat<br>Asn<br>15   | gct<br>Ala       | aca<br>Thr       | aat<br>Asn        | gga<br>Gly        | cct<br>Pro<br>20  | 201 |  |
| ggt gg<br>Gly Gl   | a aat tac<br>y Asn Tyr             | agt g<br>Ser V<br>25  | tg aca<br>al Thr       | tgg<br>Trp       | aga<br>Arg        | gat<br>Asp<br>30 | aca<br>Thr         | ggg<br>Gly       | aac<br>Asn       | ttt<br>Phe        | gtt<br>Val<br>35  | gtc<br>Val        | 249 |  |
| ggt aa<br>Gly Ly   | a ggc tgg<br>s Gly Trp<br>40       | gaa a<br>Glu I        | tc ggt<br>le Gly       | tca<br>Ser       | cca<br>Pro<br>45  | aat<br>Asn       | cga<br>Arg         | acg<br>Thr       | atc<br>Ile       | cat<br>His<br>50  | tac<br>Tyr        | aat<br>Asn        | 297 |  |
| gct gg<br>Ala Gl   | nt gtc tgg<br>y Val Trp<br>55      | gaa c<br>Glu P        | cg tct<br>ro Ser       | gga<br>Gly<br>60 | aat<br>Asn        | gga<br>Gly       | tat<br>Tyr         | ttg<br>Leu       | act<br>Thr<br>65 | ctc<br>Leu        | tat<br>Tyr        | ggg<br>Gly        | 345 |  |
| tgg ac<br>Trp Th<br>70   | a agg aat<br>ir Arg Asn<br>)       | cag c<br>Gln L        | tc ata<br>eu Ile<br>75 | gaa<br>Glu       | tat<br>Tyr        | tat<br>Tyr       | gtc<br>Val         | gtt<br>Val<br>80 | gat<br>Asp       | aat<br>Asn        | tgg<br>Trp        | gga<br>Gly        | 393 |  |
| act ta<br>Thr Ty<br>85   | ıc aga cct<br>r Arg Pro            | act g<br>Thr G<br>9   | ly Thr                 | cat<br>His       | cga<br>Arg        | ggc<br>Gly       | acc<br>Thr<br>95   | gtt<br>Val       | gtc<br>Val       | agt<br>Ser        | gat<br>Asp        | ggg<br>Gly<br>100 | 441 |  |
| gga ac<br>Gly Th   | a tat gac<br>ır Tyr Asp            | atc t<br>Ile T<br>105 | at acg<br>yr Thr       | act<br>Thr       | atg<br>Met        | Arg<br>110       | tac<br>Tyr<br>ge 1 | aat<br>Asn       | gca<br>Ala       | cct<br>Pro        | tcc<br>ser<br>115 | att<br>Ile        | 489 |  |

WO 2005/077191 PCT/DK2005/000033

| gat<br>Asp  | ggg<br>Gly        | aca<br>Thr        | caa<br>Gln<br>120 | acg<br>Thr | ttc<br>Phe        | caa<br>Gln        | cag<br>Gln        | ttc<br>Phe<br>125 | tgg<br>Trp | agt<br>Ser        | gtg<br>Val        | agg<br>Arg        | caa<br>Gln<br>130 | tcg<br>Ser | aag<br>Lys        | 537 |
|---|-------------------|-------------------|-------------------|------------|-------------------|-------------------|-------------------|-------------------|------------|-------------------|-------------------|-------------------|-------------------|------------|-------------------|-----|
| aga<br>Arg  | ccg<br>Pro        | act<br>Thr<br>135 | gga<br>Gly        | aat<br>Asn | aac<br>Asn        | gtt<br>Val        | agc<br>ser<br>140 | att<br>Ile        | acg<br>Thr | ttt<br>Phe        | agc<br>Ser        | aac<br>Asn<br>145 | cac<br>His        | gtg<br>Val | aat<br>Asn        | 585 |
| gcg<br>Ala  | tgg<br>Trp<br>150 | aga<br>Arg        | aat<br>Asn        | gca<br>Ala | gga<br>Gly        | atg<br>Met<br>155 | aat<br>Asn        | ctg<br>Leu        | gga<br>Gly | agt<br>Ser        | agt<br>Ser<br>160 | tgg<br>Trp        | tct<br>Ser        | tac<br>Tyr | cag<br>Gln        | 633 |
| gta<br>Val<br>165   | tta<br>Leu        | gca<br>Ala        | aca<br>Thr        | gaa<br>Glu | ggc<br>Gly<br>170 | tat<br>Tyr        | caa<br>Gln        | agt<br>Ser        | agc<br>Ser | ggg<br>Gly<br>175 | aga<br>Arg        | tcg<br>Ser        | aat<br>Asn        | gta<br>Val | acg<br>Thr<br>180 | 681 |
| gtt tgg tagaacgaga aagacggaat taactttctg aatatttaaa aacaaatcta 7<br>Val Trp |                   |                   |                   |            |                   |                   |                   |                   |            |                   |                   |                   | 737               |            |                   |     |
| ttgttgtgac gaacttaaga tttactcatt aagaagaatg aagc                            |                   |                   |                   |            |                   |                   |                   |                   |            |                   |                   |                   | 781               |            |                   |     |
| <210> 2<br><211> 210<br><212> PRT<br><213> Bacillus halodurans              |                   |                   |                   |            |                   |                   |                   |                   |            |                   |                   |                   |                   |            |                   |     |
| <400  | )> 2              | 2                 |                   |            |                   |                   |                   |                   |            |                   |                   |                   |                   |            |                   |     |
| Met   | Phe               | Lys               | Phe<br>-25        | Val        | Thr               | Lys               | ۷al               | Leu<br>-20        | Thr        | ٧a٦               | ٧a٦               | Ile               | Ala<br>-15        | Ala        | Thr               |     |
| Ile   | Ser               | Phe<br>-10        | Cys               | Leu        | Ser               | Ala               | Val<br>-5         | Pro               | Ala        | Ser               | Ala<br>-1         | Asn<br>1          | Thr               | Tyr        | Trp               |     |
| Gln<br>5  | Tyr               | Trp               | Thr               | Asp        | Gly<br>10         | Gly               | Gly               | Thr               | Val        | Asn<br>15         | Ala               | Thr               | Asn               | Gly        | Pro<br>20         |     |
| Gly   | Gly               | Asn               | Tyr               | Ser<br>25  | Val               | Thr               | Trp               | Arg               | Asp<br>30  | Thr               | Gly               | Asn               | Phe               | Val<br>35  | Val               |     |
| Gly   | Lys               | Glу               | Trp<br>40         | Glu        | IJе               | Glу               | Ser               | Pro<br>45         | Asn        | Arg               | Thr               | Ile               | His<br>50         | Tyr        | Asn               |     |
| Ala   | Gly               | Val<br>55         | Trp               | Glu        | Pro               | Ser               | Gly<br>60         | Asn               | Gly        | Tyr               | Leu               | Thr<br>65         | Leu               | Tyr        | Gly               |     |
| Trp   | Thr<br>70         | Arg               | Asn               | Gln        | Leu               | 11e<br>75         | Glu               | Tyr               | Tyr        | ٧a٦               | ∨a1<br>80         | Asp               | Asn               | Trp        | Gly               |     |
| Thr<br>85   | Tyr               | Arg               | Pro               | Thr        | G]у<br>90         | Thr               | ніѕ               | Arg               | Gly        | Thr<br>95         | ۷a٦               | ۷a٦               | Ser               | Asp        | Gly<br>100        |     |
| Gly   | Thr               | Tyr               | Asp               | Ile<br>105 |                   | Thr               | Thr               | Met               | Arg<br>110 |                   | Asn               | ΑΊа               | Pro               | ser<br>115 | Ile               |     |
| Asp   | Gly               | Thr               | Gln               | Thr        | Phe               | Gln               | Gln               | Phe               |            | Ser<br>ige 2      |                   | Arg               | Gln               | Ser        | Lys               |     |

WO 2005/077191 PCT/DK2005/000033

130 125 120

Arg Pro Thr Gly Asn Asn Val Ser Ile Thr Phe Ser Asn His Val Asn 140

Ala Trp Arg Asn Ala Gly Met Asn Leu Gly Ser Ser Trp Ser Tyr Gln 150 160

Val Leu Ala Thr Glu Gly Tyr Gln Ser Ser Gly Arg Ser Asn Val Thr 165 170 180

Val Trp

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(19)..(47) B. halodurans <222> <223>

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47

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source

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gcgttgagac gcgcggccgc cattcttctt aatgagtaaa tcttaagttc g

51